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 GCTGAAGCTGCTGTACAATGAAGTCCCTGTCTGGGTCCCTACCACAAGAAGTCGGCTGTAAGTGCCTTCAGTGAGGGCAGTGTC
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 GGGTTCCCTAGCTGAGTACCTCTTCCAGACTCCAACGACCCGTGCCAGGGATGTTTATGTGCAAGACTGGACGGTGCATCCGAAA
 GGAACCTGCGCTGCGACGGCTTGGCAGACTGCCCGGATTATAGTGATGAGCGTTACTGCCGATGCAATGCCACCCACAGTTTACGT
 GCAAAAACAGTTCTGCAAGCCCTCTTCTGGGTCTGTGACAGTGTCAACGACTGTGGGGACGGAAGTGACGAGGAGGGCTGCAGC
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 CTTGTGTGGGGCCTCGCTCATCTCTCTGACTGGAAGCAGCAGCTGGGTATAGCAGCATGGACAGACAGCCGACCAAAACACCCACAG
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MGSNRGRKAGGGSQDFGAGLKYNSRLENMNGFEEGVFLPANNA
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 DSCQGDSSGGLSSAEKDGRMFQAGVVSWECECAQRNKPVGVTPLPVVRDWIKEHTGV
 (SEQ ID NO:2)

FIGURE 1

underlined = deleted in targeting construct

[] = sequence flanking Neo insert in targeting construct

CATGGTAGACGGCTGCCCCGAGGGACCACGCGTCTGAGACCGGCGATCGGACCGCCAAAA
CCATGGGTAGCAATCGGGGCCGCAAGGCCGGAGGGGGCTCTCAGGACTTCGGCGCGGGAC
TCAAGTACAACCTCCCGGCTAGAGAACATGAATGGCTTTGAGGAGGGTGTGGAGTTCTTGC
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CCTACAACCTGACTTTCTCTCTCCAGAACGTCTTCTTGTACGCTGATAACCAATA
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CTGCTGGCAAGGCCATCTGGGTACAGGCTGGGGGCACACAAAAGAGGGAG [GTACCGGA
GCGCTGATCCTGCAGAAGGGTGTGATCCGTGTATCAACCAGACCACCTGTGAGGACCTC
ATGCCGACAGATCACCCACGAATGATGTGTGTGGGTTTCTCAGTGGGGGTGTGGAC
TCCTGC] CAGGGTGA CTCTGGTGGCCCCCTTGTCAAGCGCGGAGAAAG [ATGGGCGAATGT
TCCAGGCTGGTGTGGTGTGAGCTGGGGTGAAGGCTGCGCTCAGAGGAACAAGCCAGGCGTGT
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GACAGACAGCCGACCACAAACACCCACAGGGATGCCCCGACATGCACACCTGGATACAGGA
GAGGAACACTGACGACATTTATGCTGTGGCCTCCCCCCCCAACACAACCCAGACTGTGA
ACTGCATCCTTAGGACTCAGAGTTCTTCCAAAGTGGGACCCCTCAAGAGTTGGAGAGAG
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GCCCTGAGCTGGGTGAAGATGATGCTGTCCCGAGAGCTGCTTCCAAGTGTCTTGTGAGCT
CCCGGAGCCCTATGGGAGGAGGGGCTCAGGGTCACTCTTTTCAGGAAGCGCCAGCCCTA
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GTATTTGAGAGTAAACATTTTATTTCTTTTAAAAA

FIGURE 2A

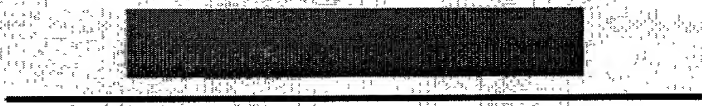
Gene Sequence Structure
*

2466 bp

Sequence Deleted

2505 bp

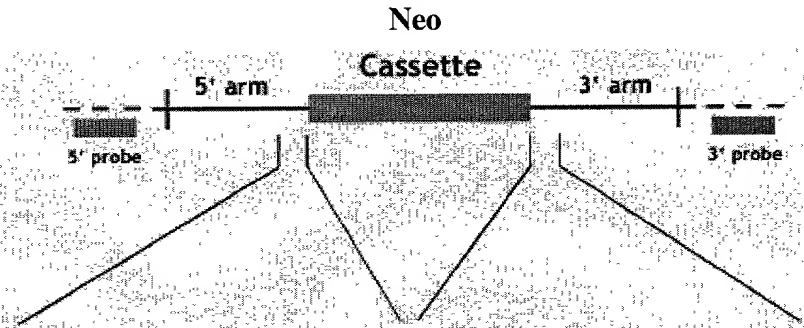
Size of full-length
cDNA: 3106 bp



Targeting Vector* (genomic sequence)

Construct Number: 2035

Arm Length:
5': 3.8 kb
3': 1 kb



<p>5' >TTCCCATTTGAGACTGGCTTA CCCCGGAAGCTGCCTGCCTCAGTC TCCCGCTTCCTGTCTCCCCAGGTA CCGGAGCGCTGATCCTGCAGAAGG GTGAGATCCGTGTCATCAACCAGA CCACCTGTGAGGACCTCATGCCGC AGCAGATCACCCACGAATGATGT GTGTGGGTTTCCTCAGTGGGGGTG TGGACTCCTGC<3' (SEQ ID NO:2)</p>	<p>5' >ATGGGCGAATGTTCCAGGCTG GTGTGGTGGGCTGGGGTGAAGGCT GCGCTCAGAGGAACAAGCCAGGCG TGTACACAAGGCTCCCTGTAGTTC GGGACTGGATCAAAGAGCACACTG GGGTATAGCAGCATGGACAGACAG CCGACCACAAACACCCACAGGGAT GCCCCGACATGCACACCTGGATACA GGAGAGGGACA<3' (SEQ ID NO:3)</p>
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FIGURE 2B